*On page 4, line 25, please replace with:*

--- Target No. 4: Residues 70-77 PDGQSTAK (**SEQ. ID. NO. 4**)---

*On page 12, lines 29-32, please replace with:*

---The VH and VL genes are liked by an artificial linker (GGGGS)$_3$ (**SEQ.ID.NO. 5**) to form

scFv fragment [VH-(GGGGS)$_3$.VL] as in Batra, et al., 1990, and the scFv fragment is subcloned

into a pBlueScript vector (Stratagene, CA).---

*On page 13, lines 11-13, please replace with:*

---scFv fragments are linked with the pentamerization domain of COMP via a hinge region

(**SEQ.ID.NO. 6**) to give rise to the following fragment: scFv-hinge [(PQ)$_2$ PK(PQ)$_4$

PKPQPK(PE)$_2$]- Pentamerization domain (COMP aa 28-72).---

*On page 18, lines 1-3, please replace with:*

---Sequence carried by the selected phage are then determined using the Sequence kit (United

States Biochemical) with the primer 5'-CCCTCATAGTTAAGCGTAACG-3'(**SEQ. ID. NO.**

**15**) (Koivunen, E., et al. (1993) J. Bio. Chem 268:20205-20210).---

*On page 18, line 18, please replace with:*

---Target No. 1: VCRHR (**SEQ. ID. NO. 7**) & GHRCL (**SEQ. ID. NO. 8**)---

*On page 18, line 19, please replace with:*

---Target No. 2: LGLVTG (**SEQ. ID. NO. 9**) & RTLVGF (**SEQ. ID. NO. 10**)---